|  |  |  |  |  |
| --- | --- | --- | --- | --- |
| **Name** | **Sequence** | **Length** | **Tm** | **GC%** |
| Parvo4\_F | GCTACAGGAGCAGAAGACTAAC | 22 | 62 | 50 |
| Parvo4\_P | AGGTAAGCAATCATGTCTGCTGCTGA | 26 | 68 | 46 |
| Parvo4\_R | TTGCCACCTGGACGATAAG | 19 | 62 | 52 |

>Parvo4\_F

GCTACAGGAGCAGAAGACTAAC

>Parvo4\_P

AGGTAAGCAATCATGTCTGCTGCTGA

>Parvo4\_R

TTGCCACCTGGACGATAAG

CTTATCGTCCAGGTGGCAA

**Amplicon 88 bp**

GCTACAGGAGCAGAAGACTAACAGGCCTCCTTTATTTTTCCAGGTAAGCAATCATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA

>NC\_007018.1 Human parvovirus 4 G1, complete genome

GCTACAGGAGCAGAAGACTAACAAGCTTCATTTATTTTTCCAGGTAAGCAA**A**CATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA

>KM390024.1 Human parvovirus 4 isolate T609, complete genome

GCTACAGGAGCAGAAGACTAACAGGCCTCCTTTATTTTTCCAGGTAAGCAATCATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA

>Tetraparvo\_CHRF\_BD0003

GCTACAGGAGCAGAAGACTAACAGGCCTCCTTTATTTTTCCAGGTAAGCAATCATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA

>CHRF\_RNA\_0012\_S12\_tetraparvovirus\_NODE\_1\_length\_5407\_cov\_103.839895\_g0\_i0

GCTACAGGAGCAGAAGACTAACAGGCCTCCTTTATTTTTCCAGGTAAGCAATCATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA

>CHRF\_RNA\_0015\_S15\_tetraparvovirus\_NODE\_1\_length\_5134\_cov\_44.556609\_g0\_i0

GCTACAGGAGCAGAAGACTAACAAGCTTCATTTATTTTTCCAGGTAAGCAA**A**CATGTCTGCTGCTGATGCTTATCGTCCAGGTGGCAA